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U.S. DEPARTMENT OF COMMERCE PATENT AND TRADEMARK OFFICE

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INFORMATION DISCLOSURE STATEMENT BY APPLICANT

(USE SEVERAL SHEETS IF NECESSARY)

APPLICANT
Dumas Milne Edwarderet al
FILING DATE

GROUP Unknown

U.S. PATENT DOCUMENTS							
EXAMINER INITIAL	DOCUMENT NUMBER	DATE	NAME	CLASS SUBCLASS		FILING DATE (IF APPROPRIATE)	
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FOREIGN PATENT DOCUMENTS							
EXAMINER	DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUBCLASS	TRANSLATION	
INITIAL						YES	NO
ó	WO 96/34981	07/1-1798° 07.11.96	PCPWO				X
d	EP 625 572	44/23/94 28.11.94	epo e?		_		
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0 3	EXAMINER INITIAL	"" (UTHER DOCUMENTS (INCLUDING AUTHOR, THEF, DATE PERTINENT PAGES, ETC.)					
-	∞	Adams, et al., NATURE, "Initial Assessment of Human Gene Diversity and Expression Patterns Based Upon 83 Million Nucleotides of cDNA Sequence", Vol. 377 Supp., pp. 1-174, September 28, 1995 99. 3 -17					
loβ	6	Caminci, et al., GENOMICS, "High-Efficiency Full-Length cDNA Cloning by Biotinylated CAP Trapper", pp. 327-336 (1998) Hillier, et al., GENOME RESEARCH, "Generation and Analysis of 280,000 Human Expressed Sequence Tags", pp. 807-828-(1996)- Gregom 2 24. 6:807-828 (1906)					
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3	4	Kato, et al., GENE, "Construction of a Human Full-Length cDNA Bank", pp. 243-250 (1984) Gene 150: 243-250 (1964)					
	~	Nomura, et al., DNA RESEARCH, "Prediction of the Coding Sequences of Unidentified Human Genes. I. The Coding Sequences of 40 New Genes (KIAA0001-KIAA0040) Deduced by Analysis of Randomly Sampled cDNA Clones from Human Immature Myeloid Cell Line KG-1", Vol. 1, pp. 27-35 (1994)					
	<i>←</i>	Gunnar von Heijne, NUCLEIC ACIDS RESEARCH, "A New Method for Predicting Signal Sequence Cleavage Sites", Vol. 14, Number 11 (1986)					

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EXAMINER	a lim	DATE CONSIDERED	6/13/13	

*EXAMINER: INITIAL IF CITATION CONSIDERED, WHETHER OR NOT CITATION IS IN CONFORMANCE WITH MPEP 609; DRAW LINE THROUGH CITATION IF NOT IN CONFORMANCE AND NOT CONSIDERED, INCLUDE COPY OF THIS FORM WITH NEXT COMMUNICATION TO APPLICANT.

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